Sequence pair distances of ExhibitC.meg ClustalW (Slow/Accurate, Gonnet) Monday, June 02, 2003 9:34 AM

Percent Identity

|    | hOP1 | hOP2 | hBMP5 | hBMP6 | 60A         | hBMP2         | hBMP4            | hGDF1                   | mBMP6(Vgr1)             | zVg1                                | Орр                                       | hBMP3   |    |
|----|------|------|-------|-------|-------------|---------------|------------------|-------------------------|-------------------------|-------------------------------------|---|---|----|
|    | -    | 2    | က     | 4     | 5           | 9             | 7                | 8                       | 6                       | 10                                  | 1   | 12  |    |
| 12 | 18.6 | 19.7 | 17.8  | 21.0  | 12.7        | 26.0          | 22.3             | 19.6                    | 15.9                    | 20.6                                | 17.2                                      |   | 12 |
| 11 | 25.1 | 26.6 | 22.9  | 23.2  | 20.4        | 38.6          | 40.7             | 19.4                    | 22.9                    | 23.1                                |   | 264.0   | 11 |
| 10 | 29.9 | 23.9 | 24.2  | 30.1  | 19.7        | 31.3          | 29.9             | 34.4                    | 31.0                    |                                     | 179.2                                     | 243.0   | 10 |
| 6  | 61.0 | 50.7 | 60.1  | 91.4  | 32.1        | 32.8          | 30.9             | 22.0                    |                         | 158.9                               | 173.1 196.7 104.1 102.0 214.0 173.3 179.2 | 260.0 285.0 261.0 237.0 255.0 265.0 243.0 264.0 | 6  |
| 8  | 16.9 | 22.3 | 18.8  | 21.2  | 12.6        | 21.2          | 21.5             |                         | 209.0                   | 162.5 195.1 141.4 158.3 123.3 158.9 | 214.0                                     | 255.0   | 8  |
| 7  | 28.2 | 30.6 | 29.7  | 29.7  | 24.0        | 62.1          |                  | 194.5                   | 140.5 129.0 136.6 209.0 | 158.3                               | 102.0                                     | 237.0   | 7  |
| 9  | 30.8 | 27.3 | 29.8  | 31.6  | 27.0        |               | 48.2             | 209.0 226.0 207.0 194.5 | 129.0                   | 141.4                               | 104.1                                     | 261.0   | 9  |
| 5  | 33.4 | 30.6 | 31.3  | 31.2  |             | 175.7         | 137.4 189.1 48.2 | 226.0                   | 140.5                   | 195.1                               | 196.7                                     | 285.0   | 2  |
| 4  | 61.7 | 50.2 | 59.3  |       | 136.8       | 3 131.9 175.7 | 137.4            | 209.0                   | 8.8                     | 162.5                               | 173.1                                     | 260.0   | 4  |
| က  | 64.3 | 50.7 |       | 55.7  | 7           | 146.          | 147.5            | 0                       | 52.5                    | 188.1                               | 165.2                                     | _   | 3  |
| 2  | 49.8 |      | 72.4  | 6.92  | 144.0 136.2 | 162.3         | 150.3            | 193.6 219.              | 6.77                    | 165.1                               | 172.6                                     | 285.0 271.0 276.                                | 2  |
| -  |      | 75.0 | 42.2  | 51.3  | 132.3       | 142.3         | 148.4            | 199.0                   | 51.5                    | 166.3                               | 165.5                                     | 285.0   | -  |
|    |      | 2    | 8     | 4     | 2           | 9             | _                | 8                       | 6                       | 0                                   | _   | 2   |    |

Divergence

Alignment Workspace of ExhibitC.meg ClustalW (Slow/Accurate, Gonnet) Monday, June 02, 2003 9:34 AM

| 1   | 100 | 37<br>27<br>62<br>42<br>46<br>59<br>59<br>59<br>59  | 200<br>200<br>200<br>200<br>51<br>48<br>49<br>61<br>61<br>49<br>53<br>25<br>25   |
|---|-----|---|--|
|   | 06  |   | 130   140   150   160   170   180   190   20   |
|   | 8,0 | PGRTEQPPPSPQ  | 180  |
|   | 7,0 | PGR   | 170  |
|   | 9   | FSL   | The control of the co |
|   | 50  | ADFSL GG  | 150  |
|   | 40  | IMAPLFTLRSALADLMSCWVLVGYAKGGPPLRPPLPAAAAAAGGLGWVLIMEVATTPPA-AGLVPELGRRKFAAASAGLVPELGRRKKVAETQGLALLLPSLPLTRAPPPLRPPLPVAAAAGGLRACLLTISLCSPALRACLLTISLCSPATRACLLTISLCSPA | 140  |
| Ω (Ω)   | 30  | - 1   |  |
| ;   | 20  | ISFVA GE GELCSCGB- VSIG VILIGGA VILIGGA VILIGGA VILIC VILICS-CGB- VILICS-CGB VILICS-CGB CILICS-CGB CILICS-CGB CILICS-CGB CILICS-CGB COSILAGG  | 120  |
| W CT CUDIT NT ACT AT A COLUMN TO A COLUMN | 10  | MHVRSLRAAAPHSFVAMILIVFLLKGIVGF MPGICRRAQWICWWWGILCSCCGP MSGIRNTSEAVAVLASIG MYAGTRCTLALLLRQVILGGA MI PENRMI MVVLCQVILGGAS MI PENRMI MVLCQVILGGAS MPPPQQGPCGHHILLL MPGICRRAQWICWWWGILCS-CGP MRGMILLLAVLATFQTIVRVASTEDI  | 110 120  |
|   |     | hOP1<br>hOP2<br>hBMP5<br>hBMP6<br>60A<br>hBMP2<br>hBMP4<br>hGDF1<br>zVg1<br>zVg1<br>hBMP3   | hOP1<br>hOP2<br>hBMP5<br>hBMP6<br>60A<br>hBMP2<br>hBMP4<br>hGDF1<br>aBMP6 (Vgr1)<br>zVg1<br>DPP  |
|   |     | hOP1<br>hOP2<br>hBMP5<br>60A<br>hBMP2<br>hBMP4<br>hGDF1<br>mBMP6<br>zVg1<br>DPP   | hOP1<br>hOP2<br>hBMP6<br>60A<br>hBMP2<br>hBMP2<br>hGDF1<br>mBMP6<br>zVg1<br>DPP  |

Alignment Workspace of ExhibitC.meg ClustalW (Slow/Accurate, Gonnet) Monday, June 02, 2003 9:34 AM

| 墓   | SAPLFMLDLYNAMSGE<br>280 290 300 | SAPMFMLDLYNAMAVE 94SAPLFMLDLYHAMGED 88SAPLFMLDLYNAMTNE 92 PGRLKSAPLFMLDLYNALSADNDEDGA 155KSAPKFLLDVYHRITAE 107AVVPPYMLDLYRRISGO 94AVIPDYMRDLYRLOSGE 94PVPPVMWRLFRRDDQ 73 PGRLKSAPLFMLDLYNALSNDDEEDGA 156AVPSQMWKLFRQASKQ 63AVPSQMWKLFKQASKQ 63 | 350 370 380 400  GENERALIDADIVMSEVNILVELDKDESHSHRHEKEFKEDLSC  350 370 380 390 400  GENERALIDADIVMSEVNILVEHDKEFFHPRYHHREFREDLTC  380 390 400  GENERALIDADIVMSEVNILVEHDKEFFHPRYHHREFREDLTC  380 390 400  GENERALIDADIVMSEVNILVEHDKEFFHPRYHHREFREDLTC  380 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 390 390 400  380 30 30  380 30 400  380 400  380 400  380 400  380 400  380 400  380 400  380 400  380 400  380 400  380 400  4 |
|---|---------------------------------|--|--|
|   | <u>1.PHRPRP</u> 250 260 270     | RREWORETLANLGLEPHREPRHL  | 35(<br>35(<br>35(<br>35(<br>35(<br>35(<br>35(<br>35(<br>35(<br>35(   |
| 部のない。  「本のない」  「本のない」 | 210 220 230 240                 | IKDKLÆPDPS   | BGERCAAHSL  310 320 340  -BG 330 340  -BG 330 340  -BG 320 350 340  -BG  |
|   |                                 | hOP1<br>hOP2<br>hBMP5<br>hBMP6<br>60A<br>hBMP2<br>hBMP4<br>hGDF1<br>mBMP6 (Vgr1)<br>zVg1<br>Dpp  | hOP1<br>hOP2<br>hBMP5<br>hBMP6<br>60A<br>hBMP2<br>hBMP4<br>hGDF1<br>mBMP6 (Vgr1)<br>zVg1<br>DPP  |

Alignment Workspace of ExhibitC.meg ClustalW (Slow/Accurate, Gonnet) Monday, June 02, 2003 9:34 AM

| MANUAL MATERIAL MATER | I PEGEAVTRAEFRIYKOYIRERFD-NEITRRISVYQVLQEHLGRESDLFILDSRITMASEBGWLVFDITATSNHWVVNPRHNLG 214 I PAGEAVTRAEFRIYK-VPSIHLL-NRTIHVSWFQVVQEQSNRESDLFFILDTRKAQALDVGWLVFDITATSNHWVYNPQNNLG 26 I PEGEAVTRAEFRIYKDRSNNRFE-NEITKISTYQIIKEYTNRDADLFTLDTRKAQALDVGWLVFDITATSNHWUINPQNNLG 26 I PEGEVATRAEFRIYKDRSNNRFF-NEITKISTYQVLQEHQHRDSDLFTLDTRKAQALDVGWLVFDITATSNHWUINPQNNLG 26 I PEGEVATRAEFRIYKDRSKWLTRNREFTISTYQVLQEHQHRDSDLFTLDTRVVMASEBGWLEFDITATSNLWVVTPQHNWG 22 I PEGETTSAEL QVFREQWDALG-NNSSFHHRINIYEVMRPPAEVVPGHLITRLLDTRLVNQNASRWESFDVTRAVNRWTRGGHANHG 22 I PENER I SSAELRLFREQVDGGPD-WERGFHRINIYEVMRPPAEVVPGHLITRLLDTRLVNHNVTRWEITEDVSPAVLRWTREKQPNYG 23 VEPAERPSRARLEI RFAAAAAAAPBGGWE | LOIEVETLDGRHVNPRRAGIVGPONTETGASEVERRITRSASRR-Q-RNKSOOSOSVRPSNVSDSACKRHELY  510 520 540 560 600  510 520 540 590 600  10IEVETLDGRHVNPRRAGIVGPONTETGASEVERTIRSASRR-Q-RNKSOOSOSVRPSNVSDSACKRHELY  33( 1RILVVETEDGYSINPKLAGILGRHGPQNKQPFNVAFFRATEVHERSIRSTGSKQRSQNRSKTPRNQEALRWANVAENSSDQAAPKHELY  1RILVVETEDGRSINVKSAGILGGRAPRSQQPFVVTFFRASEVILRSVRSANKRKNQINRNKSSRQDSRNKSDQANTSEQXQARKKHELY  1QISVVTRDGRSINVKSAGILGGRAPRSQQPFVVTFFRASEVILRSVRSANKRKNQINRNKSSRQDDSRNKSTGACKKHELX  1QISVVTRDGRSINVKSAGILGGRAPRSQQPFVATTGHDGHSRASSRRRQQSRNKSSTGSQSADYNSEGXADYNTSEQXQATLX  1QISVVTRDGRSINVKSAGILGGRAPRSQQPFVATTGHDGHSRASPTLKRR |
|--|---|--|
| IPEGEAVIAAEI RIYKDAVSGAFK-1410 420   | IPEGEAVTAAEFRIYKOYIRERFDNIPAGEAVTAAEFRIYK-VPSIHLLNIPHGEAVTAAEFRIYKORSUNRFENIPEGEVTAAEFRIYKORSUNRFENVPNINYLVWAEIRIYQNANEGKWITAYIPEFITSAELQVFREQWQGPD-WIPEAISSAEIRIFREQWQGPD-WIPEGERVTAAEFRYKOCWGSFKNIPEGEVISLAQLEWKFKQDLLLIGNIPADEKLKAAELQLTRDALSQWV-ASSQLLGHLSVIWAKSHRDIWSWLSKDI  | ELOLEVETLDGRH  510  1QLSVETLDGQS  LQLSVETEDGHS  LQLCAETGDGHS  LQLCAETGDGWS  LQLSVVTRDGWS  LQLSVVTRDGWHS  LYLGAHAVNRPDRE  FVVEVAHLEEKQGVSKRH  LALEVTHLAQTRTHQCQH  LALEVTHLAQTRTHQCQH  LALEVTHLAQTRTHQCQH  LALEVTHLAQTRTHQCQH  LALEVTHLAQTRTHQCQH  LALEVTHLAQTRTHQCQH  LALEVTHLAQTRTHQCQH  LALEVTHLAGTRTHQCQH  IQLSVVTRDGLH  MQLELQWMHINGLH  ILVEVRTVRSLKPAPHHH  IRAALSIERRKKRSTGVI  |
|  | hOP1<br>hOP2<br>hBMP5<br>hBMP6<br>60A<br>hBMP2<br>hGDF1<br>mBMP6 (Vgr1)<br>zVg1<br>DPP  | hOP1<br>hOP2<br>hBMP5<br>hBMP6<br>60A<br>hBMP2<br>hBMP4<br>hGDF1<br>mBMP6 (Vgr1)<br>zVg1<br>DPP  |

Alignment Workspace of ExhibitC.meg ClustalW (Slow/Accurate, Gonnet) Monday, June 02, 2003 9:34 AM

| NSFSDIGWODWIIAPEGYAANYCHGECSFPIANAHMAATNHAIVOTIVHIMNPEK-VPKPCCAPIKISAISVLYFDDNSNVVIKKYRNMVVRACGCH 610 620 630 640 650 660 670 680 690 700 | VSFRDIGWOMI IAPBGYAAYYCEGECAFPINSYMNATNHAIVQIIJHFINPET-VPRCCAPTQIAALSVLYFDDSSNVILKKYRNMVVRAGGH 402 VSFQDIGMIDMVIAPQGYSAYYCEGECSFPLDSGMNATNHAILQSIJHIMKFNA-VPRACCAPTKIAALSVLYYDSSNVVILKKYRNMVVRAGGH 454 VSFQDIGMIDMVIAPQGYSAYYCEGECSFPLDSGMNATNHAILQSIJHIMKFNA-VPRACCAPTKIAALSVLYFDDSSNVILKKYRNMVVRAGGH 454 VSFQDIGMQDMI IAPBGYAAFYCGECSFPLNAHMNATNHAIVQTIJHIMNPEY-VPRECCAPTKIAALSVLYFDDSSNVILKKYRNMVVRAGGH 455 IDFROIGMHDMI IAPBGYGAFYCSECNFPLAAHMNATNHAIVQTIJHIMNPEY-VPRECCAPTKIAALSVLYFDDNSNVILKKYRNMVVRAGGH 455 VDFSDVGMNDMIVAPPGYAFYCHGECPFPLADHINSTNHAIVQTIJNNSVN-SK-IPKACCVPTELSALSMLYLDENDKVLKNYQDMVVBGGGR 406 VSFREVGMNDMIVAPPGYAFYCHGECPFPLADHINSTNHAIVQTIJNNSVN-SK-IPKACCVPTELSALSMLYLDENDKVVLKNYQDMVVBGGGR 376 VSFREVGMNDMIVAPPGYARYCHGECPFPLADHINSTNHAIVQTIJHIMNPEY-VPRECCAPTKIABISTIJHFTNSDNVILKKYRNMVVBGGGR 376 VSFREVGMDMI IAPKGYAANYCGGCSFPINAHMNATNHAIVQTIJHIMNPEY-VPKPCCAPTKIASISMLYTDDNSNVILKKYRNMVVBCGCR 356 VDFSDVGMDMI IAPQGYLANYCGGCSFPINAHMNATNHAILQTIJHSFDDRKG-TPQPCCAPTKIASISMLYTDDSTAVLKNYZRNMVVBCGCR 356 VDFSDVGMDMIVAPIGYDAYYCHGKCPFPIADHFNSTNHAIVQTIJNNNMNFGK-VPKACCVPTQILDSVAMLYIADQSTVVLKNYZRNMTVVBCCRR 472 VDFADIGMSEMI ISPKSFDAYYCGGCOPTRIABINGSISILFFDENKNVVLKNYYPNMTVVBCCRR 472 | 433<br>405<br>456<br>456<br>313<br>375<br>375<br>375<br>375<br>406<br>406<br>375<br>375<br>375<br>375 |
|---|---|---|
| ī   |   |   |
|   | hOP1<br>hOP2<br>hBMP5<br>hBMP6<br>60A<br>hBMP2<br>hBMP4<br>hGDF1<br>mBMP6 (Vgr1)<br>zVg1<br>DPP   | hOP1<br>hOP2<br>hBMP5<br>hBMP6<br>60A<br>hBMP2<br>hGDF1<br>mBMP6 (Vgr1)<br>zVg1<br>DPP                |

Sequence pair distances of ExhibitC.meg ClustalW (Slow/Accurate, Gonnet) Monday, June 02, 2003 9:34 AM

|                  |    | -    | 2    | ဗ    | 4    | 5                       | 9                             | 7                       | ω                                   | 6                       | 9                 | 7   | 12  |    |
|------------------|----|------|------|------|------|-------------------------|-------------------------------|-------------------------|-------------------------------------|-------------------------|-------------------|---|---|----|
|                  | 12 | 18.6 | 19.7 | 17.8 | 21.0 | 12.7                    | 26.0                          | 22.3                    | 19.6                                | 15.9                    | 20.6              | 17.2  |   | 12 |
|                  | 11 | 25.1 | 26.6 | 22.9 | 23.2 | 20.4                    | 38.6                          | 40.7                    | 19.4                                | 22.9                    | 23.1              |   | 264.0   | 11 |
|                  | 10 | 29.9 | 23.9 | 24.2 | 30.1 | 19.7                    | 31.3                          | 29.9                    | 34.4                                | 31.0                    |                   | 179.2   | 285.0 271.0 276.0 260.0 285.0 261.0 237.0 255.0 265.0 243.0 264.0 | 10 |
|                  | 6  | 61.0 | 50.7 | 60.1 | 91.4 | 32.1                    | 32.8                          | 30.9                    | 22.0                                |                         | 158.9             | 173.3   | 265.0   | 6  |
| λ                | 8  | 16.9 | 22.3 | 18.8 | 21.2 | 12.6                    | 21.2                          | 21.5                    |                                     | 209.0                   | 123.3             | 214.0   | 255.0   | æ  |
| Identit          | 7  | 28.2 | 30.6 | 29.7 | 29.7 | 24.0                    | 62.1                          |                         | 194.5                               | 136.6                   | 158.3             | 102.0   | 237.0   | 7  |
| Percent Identity | 9  | 30.8 | 27.3 | 29.8 | 31.6 | 27.0                    |                               | 48.2                    | 207.0                               | 129.0                   | 141.4             | 104.1   | 261.0   | 9  |
| م                | 5  | 33.4 | 30.6 | 31.3 | 31.2 |                         | 175.7                         | 189.1                   | 226.0                               | 140.5 129.0 136.6 209.0 | 162.5 195.1 141.4 | 172.6 165.2 173.1 196.7 104.1 102.0 214.0 173.3 179.2 | 285.0   | 2  |
|                  | 4  | 61.7 | 50.2 | 59.3 |      | 136.8                   | 131.9                         | 150.3 147.5 137.4 189.1 | 209.0                               | 8.8                     | 162.5             | 173.1   | 260.0   | 4  |
|                  | 3  | 64.3 | 50.7 |      | 55.7 | 136.2                   | 146.6                         | 147.5                   | 219.0                               | 52.5                    | 188.1             | 165.2   | 276.0   | က  |
|                  | 2  | 49.8 |      | 72.4 | 6.97 | 132.3 144.0 136.2 136.8 | 142.3 162.3 146.6 131.9 175.7 | 150.3                   | 199.0 193.6 219.0 209.0 226.0 207.0 | 6.77                    | 166.3 165.1 188.1 | 172.6   | 271.0   | 2  |
|                  | -  |      | 75.0 | 42.2 | 51.3 | 132.3                   | 142.3                         | 148.4                   | 199.0                               | 51.5                    | 166.3             | 165.5   | 285.0   | -  |
|                  |    |      |      |      |      |                         |                               |                         |                                     |                         |                   | _   | ~   |    |

hOP2 hBMP5 hBMP6 60A hBMP2 hBMP4

hOP1

mBMP6(Vgr1)

zVg1 Dpp hBMP3

Divergence